RAW SEQUENCE LISTING

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Application Serial Number: 10/521319 PCT
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RAW SEQUENCE LISTING DATE: 01/28/2005
PATENT APPLICATION: **US/10/521,319** TIME: 11:41:28

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     9 <120> TITLE OF INVENTION: Animal Models
    13 <130> FILE REFERENCE: PG4871
C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/521,319
C--> 19 <141> CURRENT FILING DATE: 2005-01-14
    23 <160> NUMBER OF SEQ ID NOS: 20
    27 <170> SOFTWARE: PatentIn version 3.1
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    33 <211> LENGTH: 2984
    35 <212> TYPE: DNA
    37 <213> ORGANISM: Homo sapiens
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Pro Ser Lys Val Ile Leu Pro Arg Gly Gly Ser Val Leu Val Thr Cys
        35                  40                  45
Ser Thr Ser Cys Asp Gln Pro Lys Leu Leu Gly Ile Glu Thr Pro Leu
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219 Pro Asp Gly Gln Ser Thr Ala Lys Thr Phe Leu Thr Val Tyr Trp Thr
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223 Pro Glu Arg Val Glu Leu Ala Pro Leu Pro Ser Trp Gln Pro Val Gly
224         115                 120                 125
227 Lys Asn Leu Thr Leu Arg Cys Gln Val Glu Gly Gly Ala Pro Arg Ala
228     130                 135                 140
231 Asn Leu Thr Val Val Leu Leu Arg Gly Glu Lys Glu Leu Lys Arg Glu
232 145                 150                 155                 160
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236                 165                 170                 175
239 Arg Asp His His Gly Ala Asn Phe Ser Cys Arg Thr Glu Leu Asp Leu
240             180                 185                 190
243 Arg Pro Gln Gly Leu Glu Leu Phe Glu Asn Thr Ser Ala Pro Tyr Gln
244         195                 200                 205
247 Leu Gln Thr Phe Val Leu Pro Ala Thr Pro Pro Gln Leu Val Ser Pro
248     210                 215                 220
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252 225                 230                 235                 240
255 Gly Leu Phe Pro Val Ser Glu Ala Gln Val His Leu Ala Leu Gly Asp
256                 245                 250                 255
259 Gln Arg Leu Asn Pro Thr Val Thr Tyr Gly Asn Asp Ser Phe Ser Ala
260             260                 265                 270
263 Lys Ala Ser Val Ser Val Thr Ala Glu Asp Glu Gly Thr Gln Arg Leu
264         275                 280                 285
267 Thr Cys Ala Val Ile Leu Gly Asn Gln Ser Gln Glu Thr Leu Gln Thr
268     290                 295                 300
271 Val Thr Ile Tyr Ser Phe Pro Ala Pro Asn Val Ile Leu Thr Lys Pro
272 305                 310                 315                 320
275 Glu Val Ser Glu Gly Thr Glu Val Thr Val Lys Cys Glu Ala His Pro
276                 325                 330                 335
279 Arg Ala Lys Val Thr Leu Asn Gly Val Pro Ala Gln Pro Leu Gly Pro
280             340                 345                 350
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284         355                 360                 365
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288     370                 375                 380
291 Asn Gln Thr Arg Glu Leu Arg Val Leu Tyr Gly Pro Arg Leu Asp Glu
292 385                 390                 395                 400
295 Arg Asp Cys Pro Gly Asn Trp Thr Trp Pro Glu Asn Ser Gln Gln Thr
296                 405                 410                 415
299 Pro Met Cys Gln Ala Trp Gly Asn Pro Leu Pro Glu Leu Lys Cys Leu
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303 Lys Asp Gly Thr Phe Pro Leu Pro Ile Gly Glu Ser Val Thr Val Thr
304         435                 440                 445
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307 Arg Asp Leu Glu Gly Thr Tyr Leu Cys Arg Ala Arg Ser Thr Gln Gly
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312 465 470 475 480
315 Ile Val Ile Ile Thr Val Val Ala Ala Ala Val Ile Met Gly Thr Ala
316 485 490 495
319 Gly Leu Ser Thr Tyr Leu Tyr Asn Arg Gln Arg Lys Ile Lys Lys Tyr
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379 Glu Pro Ala Val Gly Glu Pro Ala Glu Val Thr Thr Thr Val Leu Val
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